NCBI **Blast 2 Sequences results**

PubMed   Entrez   BLAST   OMIM   Taxonomy   Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.17 [Aug-26-2007]

Match: 1   Mismatch: -2   gap open: 5   gap extension: 2
x_dropoff: 0   expect: 10.0000 wordsize: 11   Filter ☐ View option Standard ▼
Masking character option X for protein, n for nucleotide ▼   Masking color option Black ▼
☐ Show CDS translation   Align

Sequence 1: gi|13157532|Human DNA sequence from clone RP11-428G23 on chromosome 13
Contains part of a novel gene (KIAA0916), complete sequence
Length = 401 (66400 .. 66800)

Sequence 2: lcl|65536
Length = 303 (1 .. 303)
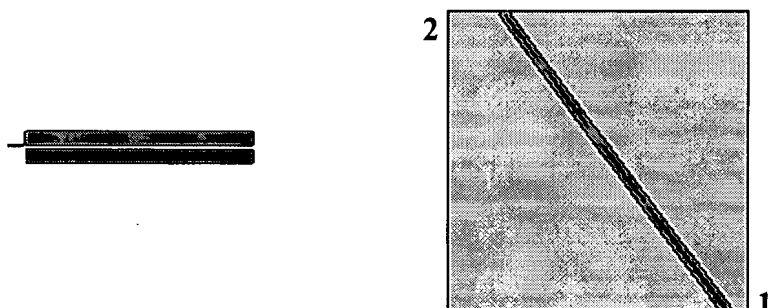
NOTE:Bitscore and expect value are calculated based on the size of the nr database.

NOTE:If protein translation is reversed, please repeat the search with reverse strand of the query sequence.


```
Score =  583 bits (303),  Expect = 2e-163
Identities = 303/303 (100%), Gaps = 0/303 (0%)
Strand=Plus/Minus

Query  66474  CTACCAACCTGTACAGCATGCTGGTGTACTGAATACTGTAGGCAACTGTAACCCATTAGT  66533
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct  303    CTACCAACCTGTACAGCATGCTGGTGTACTGAATACTGTAGGCAACTGTAACCCATTAGT  244

Query  66534  AAGTATTTCTGCATGTAAACATAGAAAAGTTATAGTCAAAGTACTTATTATAATCTTATG  66593
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct  243    AAGTATTTCTGCATGTAAACATAGAAAAGTTATAGTCAAAGTACTTATTATAATCTTATG  184

Query  66594  GGAACACCTTAGCATACGCAGTCCATCACTGACCAAAATACTGTTATACAGTGCATAACT  66653
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              | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Sbjct   183   GGAACACCTTAGCATACGCAGTCCATCACTGACCAAAATACTGTTATACAGTGCATAACT   124

Query   66654 GTGTATACACATACATACATATATAGGTATATATATATAAAATAGTGTGTCTGCATGCTT   66713
              | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Sbjct   123   GTGTATACACATACATACATATATAGGTATATATATATAAAATAGTGTGTCTGCATGCTT   64

Query   66714 ATATATCTTCACAAGAAGAAACACAAGAAGAATAAACTGTAAAACAATGACATTCATTAC   66773
              | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Sbjct   63    ATATATCTTCACAAGAAGAAACACAAGAAGAATAAACTGTAAAACAATGACATTCATTAC   4

Query   66774 CTC   66776
              | | |
Sbjct   3     CTC   1
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CPU time: 0.10 user secs.   0.04 sys. secs   0.14 total secs.